

MMU72677_1	-----	
219_HUMAN	MR-LAG-PLRIVVLVSVGVTVIVVSILLGGPGSG--FPRIQQLFTSPES	46
D87120_1_	MR-VAG-AAKLVVAVAVFLLTFYVISQVFEIKMDA--SLGNLFARSALDT	46
z219a.pep	MRPLAGGLLKVVVFVVFASLCAWYSGYLLAELIPDAPLSSAAYSIRSIGER	50
MMU72677_1	-----	
219_HUMAN	SVTAAPRARKYKCGLPQPCPEEHLAFRVVSGAANVIGPKICLEDKMLMSS	96
D87120_1_	-AARSTKPPRYKCGISKACPEKHFAFKMASGAANVVGPKICLEDNVLMMSG	95
z219a.pep	PVLKAPVPKRQKCDHWTPCPSDTYAYRLLSGGGRSKYAKICFEDNLLMGE	100
MMU72677_1	-----TGQVMKKDSFDMYSGD-PQLLLNFLTEIPDSTL	32
219_HUMAN	VKDNVGRGLNIALVNGVSGELIEARAFDMWAGD-VNDLLKFIRPLHEGTL	145
D87120_1_	VKNNVGRGINVALANGKTGEVLDTKYFDMWGGD-VAPFIEFLKAIQDGTI	144
z219a.pep	QLGNVARGINIAIVNYVTGNVTATRCFDMYEGDNSGPMTKFIQSAAPKSL	150
MMU72677_1		
219_HUMAN	VLVASYDDPGTKMNDKIKTLFSNLGSSYAKQLGFRDSWVFVGAKDLKSKS	82
D87120_1_	VFVASYDDPATKMNEETRKLFSSELGSRNAKELAFRDSWVFVGAKGVQNKs	195
z219a.pep	VLMGTYYDDGATKLNDEARRLIADLGSTSITNLGFRDNWVFCGGKGIKTKS	194
	LFMVTYYDDGSTRLNNDAKNAIEALGSKEIRNMKFRSSWVFIAAKGLELPS	200
MMU72677_1	PYEQFLKNNPET--NKYDGWPELLELEGCVPRKVM--	115
219_HUMAN	PFEQHVKNKSHS--NKYEGCPEALEMEGCIPRRSTAS	230
D87120_1_	PFEQHIKNNKDT--NKYEGWPEVVEMEGCIPQKQD--	227
z219a.pep	EIQREKINHSDAKNNRYSGWPAEIQIEGCIPKERS--	235

FIGURE